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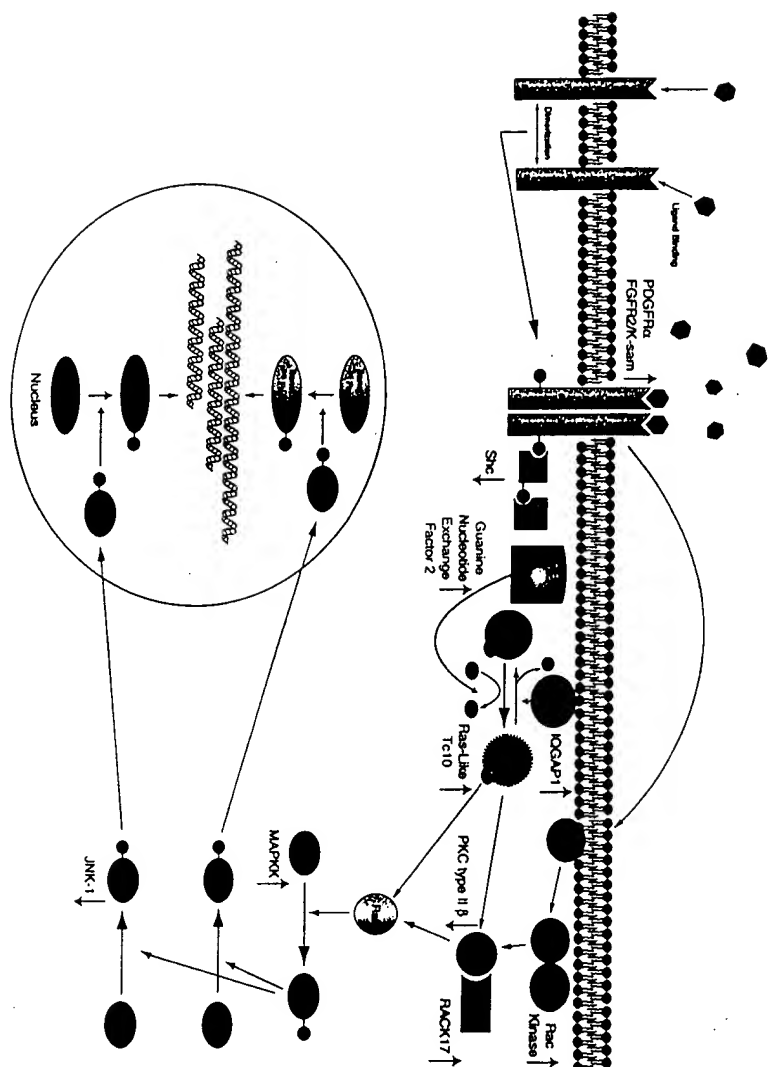
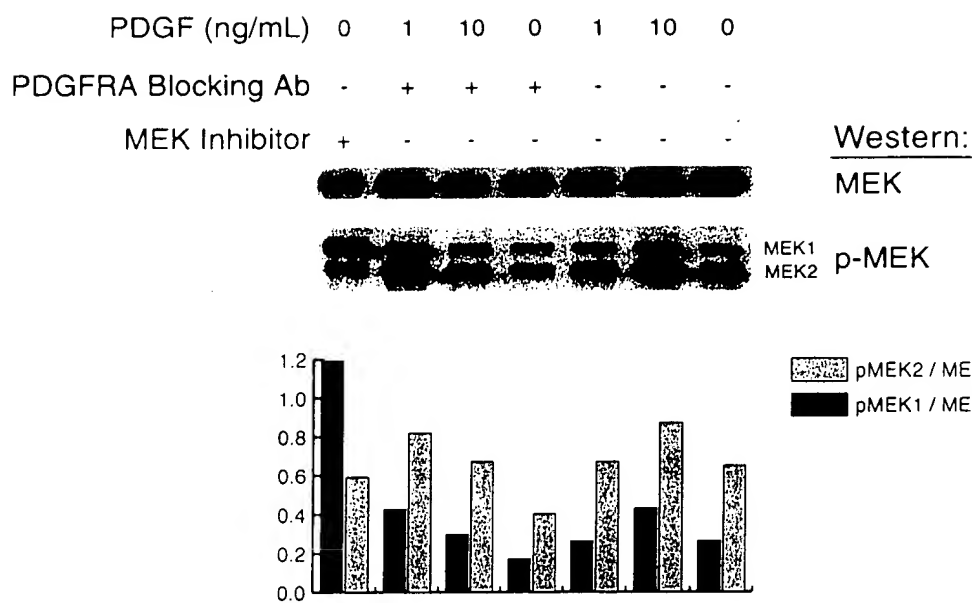


Fig. 1





a.



b.

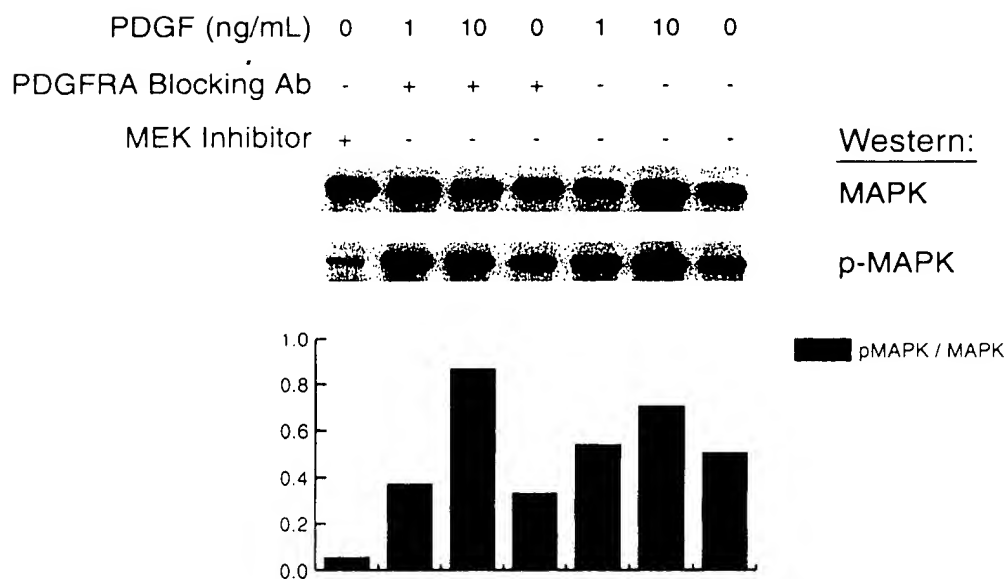


Fig. 2

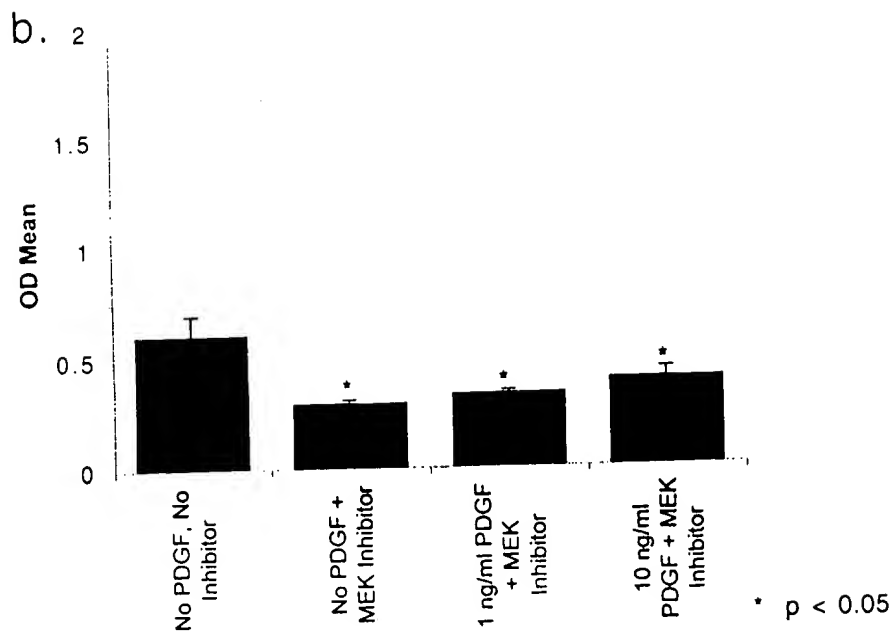
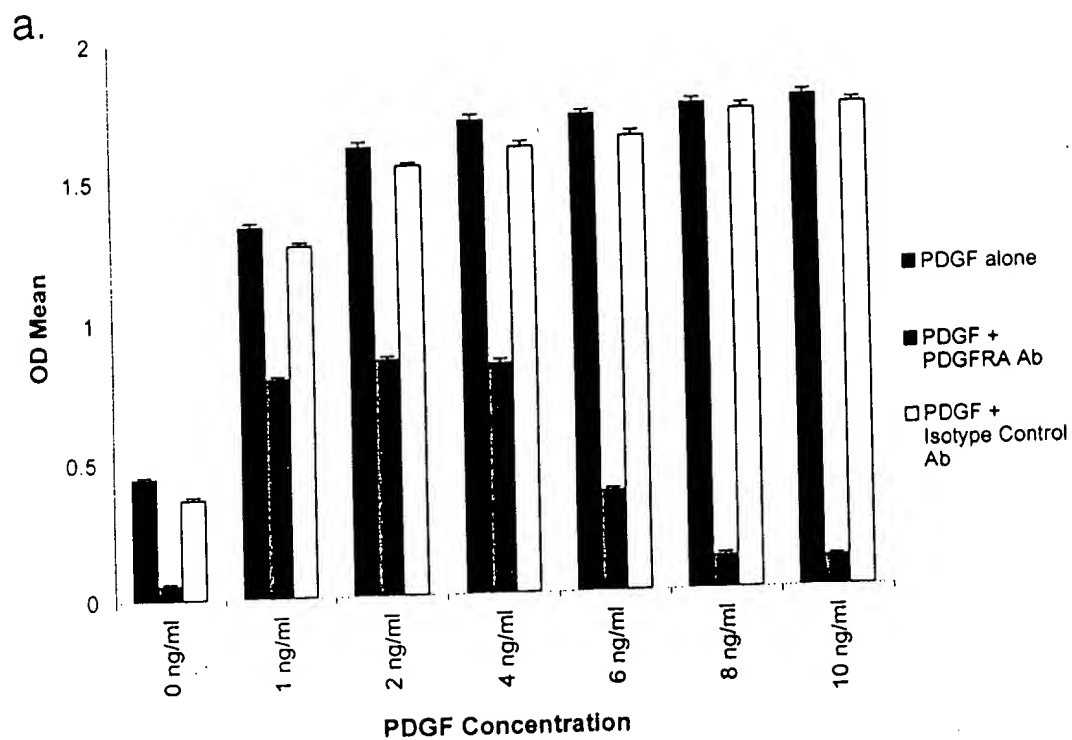
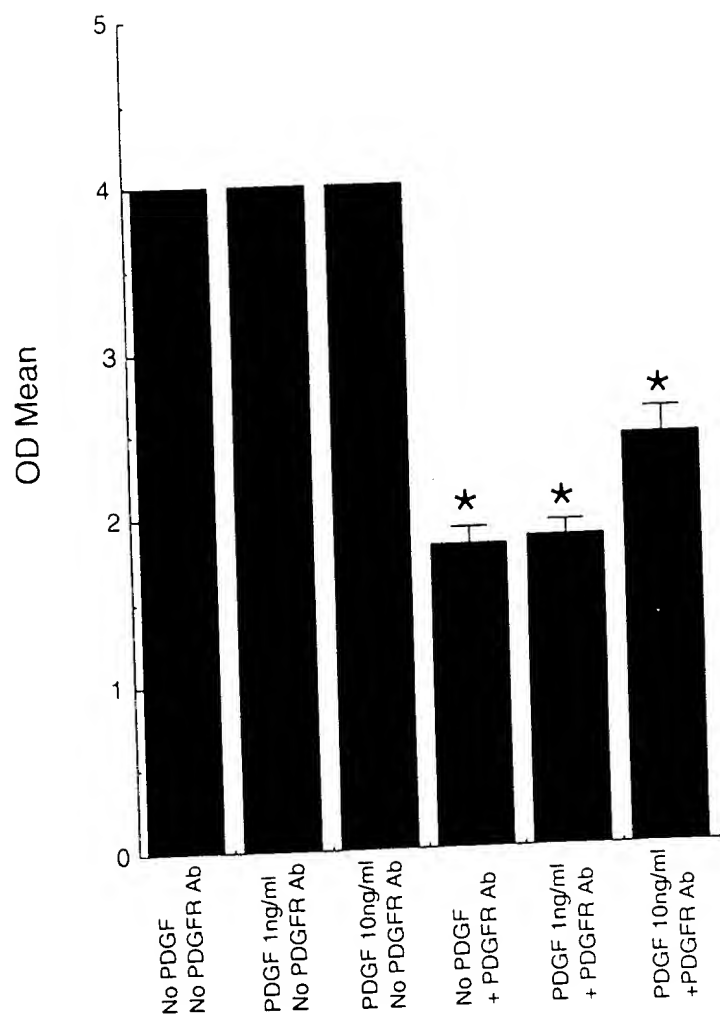


Fig. 3



\*  $p < 0.05$

Fig. 4

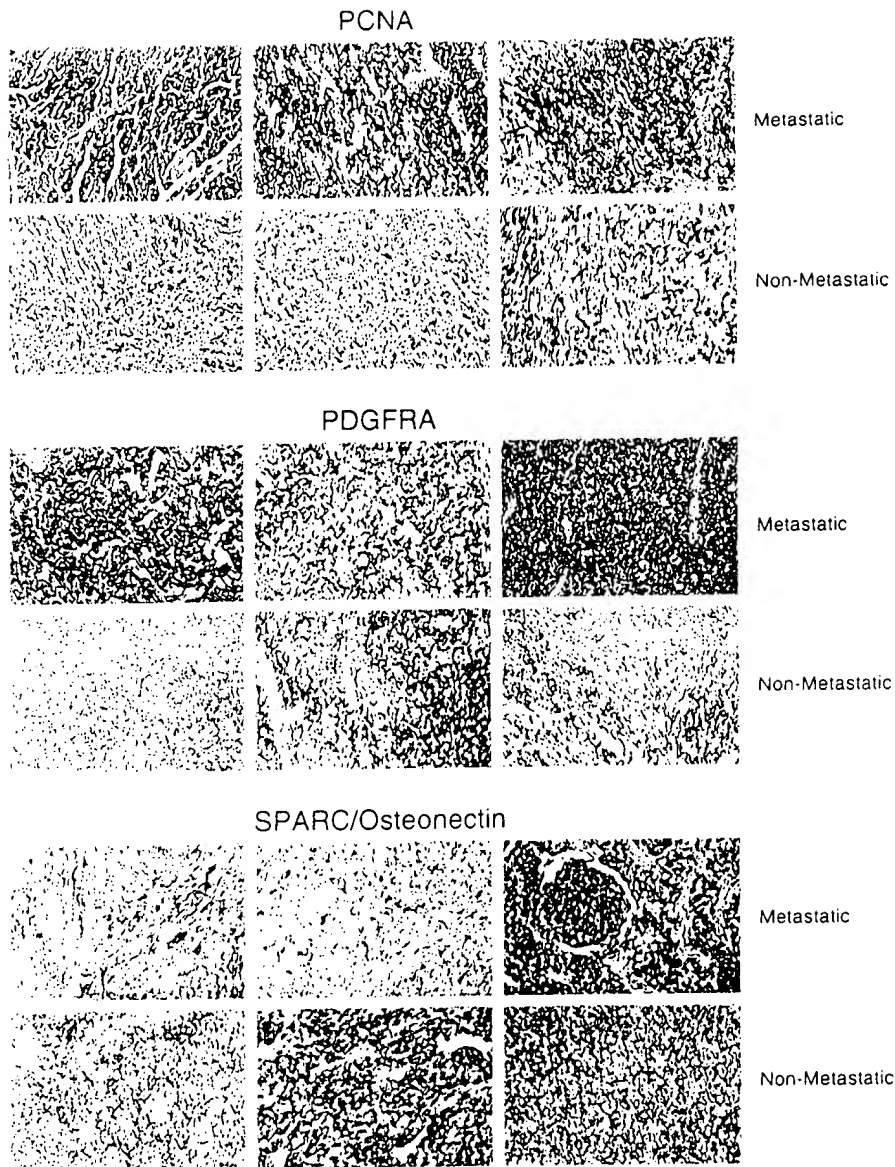


Fig. 5

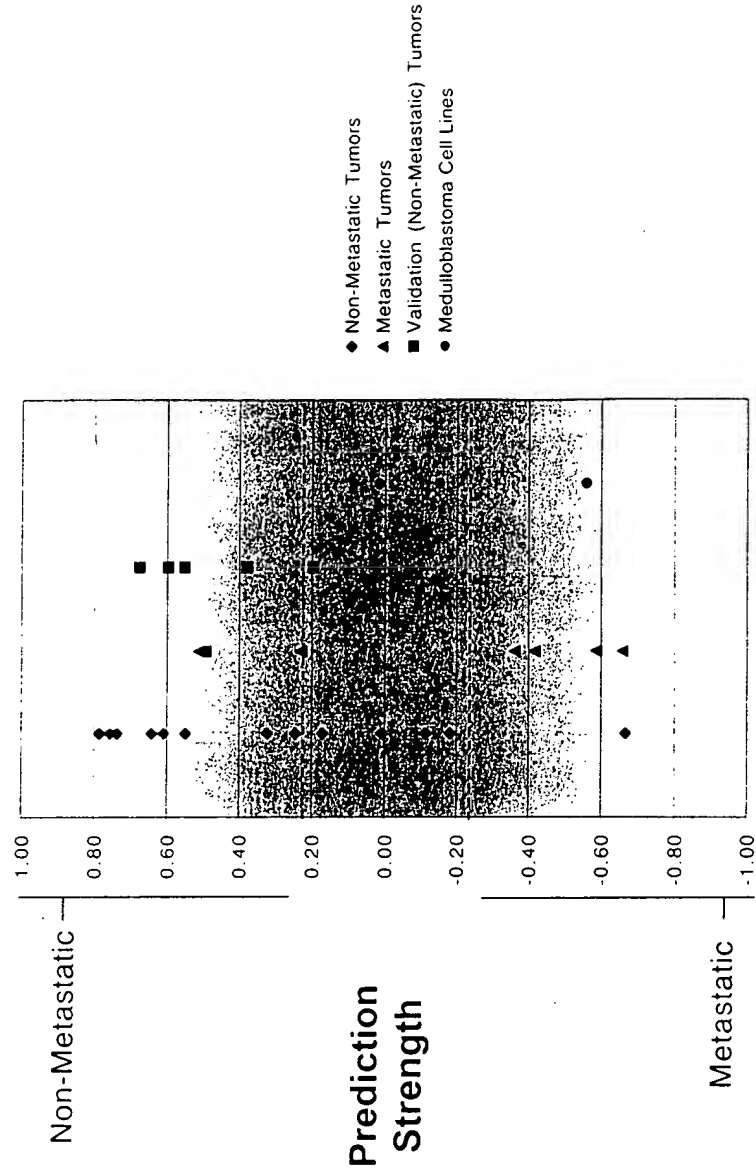


Fig. 6



Probe Set	Gene Name	Average Intensity, Non-Metastatic	Average Intensity, Metastatic	Permutational p-value	Average Fold Difference	
1937_at	Rasipoblastoma 1	3608	1966	0.004	1.80	
824_at	OTF-binding protein (RAB38)	134	23	0.006	5.83	
1811_at	Interferon (IFI-gamma)	111	38	0.007	2.88	
1848_at	Interleukin 10 (IL10)	381	171	0.007	2.22	
8043_at	e-ryb	807	237	0.018	1.71	
866_at	Integrin alpha-3 chain	1308	640	0.018	2.06	
529_at	Human dual-specificity protein phosphatase	241	71	0.024	3.42	
2070_at	Protein kinase (JNK1)	273	107	0.025	2.54	
786_at	Nedd4-like ubiquitin-protein ligase WWPE	1367	518	0.026	2.66	
1812_at	APC	67	19	0.036	4.88	
304_at	Quarantine Nucleoside Exchange Factor 2	3387	1864	0.036	2.04	
443_at	Nuclear factor I B3	239	140	0.036	1.71	
1800_at	Karstomy growth factor	322	183	0.037	1.78	
684_at	Tyrosine kinase (TQK)	3264	1818	0.037	1.80	
1447_at	MXI1	860	471	0.037	2.04	
1127_at	Epidermal growth factor receptor kinase substrate (Eps8)	657	368	0.040	2.33	
8046_at	Ribosomal protein S8 kinase 2 (RPS8K2)	718	366	0.044	1.87	
2022_at	Erg protein (steroid gene), 3' flank	128	80	0.046	2.08	
528_at	Ras protein kinase beta	468	288	0.046	1.62	
547_at	Heat shock protein 27 (HSP27)	314	11	0.047	28.22	
1116_at	THURF- NQF1-Smurf7 beta-type transcription factor homolog	162	32	0.048	5.01	
1012_at	Protein kinase C (PKC) type beta 8	130	68	0.048	1.92	
1811_at	p300/CBP-associated factor (P/CAF)	886	870	0.048	1.48	Down
788_at	p65 and p66 isoforms of N-B2	757	437	0.048	1.73	In M+
138_at	Chorionic Somatomammotropin Hormone Cα-4	62	28	0.050	2.08	
308_at	Quarantine kinase associated protein (QKAP)	18	28	0.050	1.40	
828_at	Homeobox 1.4	3362	1468	0.000	3.89	Up In M+
828_at	Gluathione S-transferase-P1s	3368	8098	0.001	2.11	
828_at	Cathepsin D (B6D)	780	1830	0.003	1.84	
1852_at	Replication protein A 140Da subunit (RPA)	184	3186	0.004	20.14	
8083_at	Tissue inhibitor of metalloproteinases (TIMP)	3366	11374	0.004	3.38	
1811_at	MAC28	185	370	0.004	1.92	
871_at	Mucin (MUC8)	217	816	0.006	2.84	
871_at	Replication protein A 140Da subunit (RPA)	4166	8688	0.007	2.08	
1718_at	Ran-Like Protein Tε10	578	1318	0.007	2.28	
1741_at	Insulin-like growth factor binding protein-2	417	3012	0.008	4.83	
841_at	Protein kinase C-binding protein RACK17	78	447	0.008	5.18	
1321_at	Tumor-associated membrane protein homolog (TMP)	90	343	0.008	3.80	
1143_at	FGF Receptor K-Sam, AL, Epitor 3	2401	3486	0.008	1.48	
1179_at	Epidermal/keratin N1-Jointlytransferrase, AL, Epitor 3	3393	8071	0.010	1.48	
708_at	Beta-tubulin gene, clone m40	80	408	0.012	5.10	
1319_at	X747840 receptor protein tyrosine kinase	368	780	0.012	2.18	
368_at	ST4 Oncolest antigen	281	748	0.013	2.67	
1001_at	Putative receptor tyrosine kinase (te)	568	847	0.013	1.52	
842_at	P1-C6468	864	1413	0.013	1.64	
1062_at	NF-κB-beta	2946	4256	0.013	1.81	
283_at	Ubiquitin-like cytochrome-c reductase core 1	332	843	0.015	1.84	
1064_at	Replication factor C, 37-42Da subunit	674	1878	0.016	2.83	
770_at	Gluathione peroxidase	513	1461	0.016	2.83	
317_at	D56686 Cysteine protease	928	1878	0.017	1.81	
1843_at	Tumor necrosis factor receptor	80	212	0.017	3.84	
1007_at	Receptor tyrosine kinase DDR	1758	2943	0.018	1.67	
1808_at	Receptor protein-tyrosine kinase (DEK8)	120	787	0.018	6.42	
928_at	Gamma-interferon-inducible protein (P-30)	787	1340	0.019	1.78	
1844_at	Bloom's syndrome protein (BLM)	443	886	0.019	1.87	
818_at	Homeobox protein (HOC7)	10	277	0.019	27.71	
1137_at	Leukemia virus receptor 2 (GLVR2)	270	519	0.019	3.03	
605_at	RNA polymerase II subunit (hRPB10)	1774	3386	0.021	1.85	
1306_at	Cytochrome P-450LTV	368	841	0.023	1.78	
1470_at	DNA polymerase delta small subunit	577	1644	0.023	2.80	
1198_at	RCC1 isoform 1-14	220	637	0.025	2.80	
814_at	Homeobox protein (HOC7)	1884	2817	0.028	2.04	
1782_at	Oncoprotein 18 (Op18)	3241	4868	0.028	1.44	
736_at	Protein Kinase H31, Camp-Dependent	11	128	0.028	18.03	
828_at	(clone 14Y8) metalloproteinase-9 (MT10)	218	548	0.028	2.57	
428_at	mRNA ligament for beta-3 integrin/alpha-5	5818	9673	0.029	1.62	
811_at	Phenoxazin, AL, Epitor 1	641	2028	0.030	3.70	
1228_at	THF-alpha converting enzyme	183	308	0.031	1.67	
1771_at	Platelet-derived growth factor receptor alpha	382	821	0.032	1.82	
1846_at	X730840c N425-H1	4198	8670	0.033	1.88	
1837_at	Ras GTPase-activating-like protein (GAP1)	368	842	0.034	1.84	
608_at	WAPKAP kinase (DpK)	67	172	0.036	2.54	
1970_at	Metastathionein-1B	3146	4404	0.036	1.40	
1378_at	FGFR2	274	881	0.037	2.48	
3068_at	M88371 Protein tyrosine kinase	346	752	0.037	2.18	
1137_at	Integrin beta-6 subunit	630	1046	0.038	1.66	
1104_at	Phosphoprotein p63	446	832	0.040	2.08	
1104_at	Heat shock protein (hsp 70)	1983	4117	0.042	2.07	
133_at	Cathepsin C	442	783	0.043	1.78	
8024_at	Lyn B	63	257	0.044	4.02	
702_at	Homeobox Protein Hps-6	344	518	0.044	1.51	
882_at	Tumor antigen (L8)	63	178	0.044	2.58	
1771_at	Maz2	803	1070	0.046	1.78	
841_at	Mutator gene (M4SH42)	721	1067	0.046	1.47	
2048_at	Alpha1(E)-catenin	2467	3607	0.046	1.42	

Fig. 7